



Epicentre-00005 seqlist (Nov).txt

SEQUENCE LISTING

<110> Davydova, Elena K.
Rothman-Denes, Lucia B.
Dahl, Gary A.
Gerdes, Svetlana Y.
Jendrisak, Jerome J.

<120> PREPARATION AND USE OF SINGLE-STRANDED TRANSCRIPTION SUBSTRATE
S
FOR SYNTHESIS OF TRANSCRIPTION PRODUCTS CORRESPONDING TO TARGE
T
SEQUENCES

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<170> PatentIn Ver. 2.1

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Thr	Asp	Ala	Thr	Pro	Glu	Gln	Lys	Val	Ala	Ala	His	Gln	Tyr	Val	Ser
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Asp	Leu	Met	Asn	Ala	Thr	Arg	Phe	Asn	Pro	Glu	Asn	Tyr	Gln	Glu	Ala
545					550					555					560
Pro	Glu	His	Ile	Arg	Asn	Ala	Val	Ala	Gly	Ser	Thr	Asp	Gln	Val	Gln
				565					570					575	
Val	Ile	Gln	Lys	Leu	Ala	Asp	Leu	Val	Asn	Thr	Leu	Asp	Glu	Ser	Asn
			580					585					590		
Pro	Gln	Ala	Leu	Met	Glu	Ala	Ala	Ser	Tyr	Met	Tyr	Asp	Ala	Val	Ser
		595					600					605			
Glu	Phe	Glu	Gln	Phe	Ile	Asn	Arg	Asp	Pro	Ala	Ala	Leu	Asp	Ser	Ile
	610					615					620				
Pro	Lys	Asp	Ser	Pro	Ala	Ile	Glu	Leu	Leu	Asn	Arg	Tyr	Thr	Asn	Leu
625					630					635					640
Thr	Ala	Asn	Ile	Gln	Asn	Thr	Pro	Lys	Val	Ile	Gly	Ala	Leu	Asn	Val
				645					650					655	
Ile	Asn	Arg	Met	Ile	Asn	Glu	Ser	Ala	Gln	Asn	Gly	Ser	Leu	Asn	Val
			660					665					670		
Thr	Glu	Glu	Ser	Ser	Pro	Gln	Glu	Met	Gln	Asn	Val	Ala	Leu	Ala	Ala
		675					680					685			
Glu	Val	Ala	Pro	Glu	Lys	Leu	Asn	Pro	Glu	Ser	Val	Asn	Val	Val	Leu

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700

Lys	His	Ala	Ala	Asp	Gly	Arg	Ile	Lys	Leu	Asn	Asn	Arg	Gln	Ile	Ala
705					710					715					720
Ala	Leu	Gln	Asn	Ala	Ala	Ala	Ile	Leu	Lys	Gly	Ala	Arg	Glu	Tyr	Asp
			725						730					735	
Ala	Glu	Ala	Ala	Arg	Leu	Gly	Leu	Arg	Pro	Gln	Asp	Ile	Val	Ser	Lys
			740					745					750		
Gln	Ile	Lys	Thr	Asp	Glu	Ser	Arg	Thr	Gln	Glu	Gly	Gln	Tyr	Ser	Ala
		755					760					765			
Leu	Gln	His	Ala	Asn	Arg	Ile	Arg	Ser	Ala	Tyr	Asn	Ser	Gly	Asn	Phe
	770					775					780				
Glu	Leu	Ala	Ser	Ala	Tyr	Leu	Asn	Asp	Phe	Met	Gln	Phe	Ala	Gln	His
785					790					795					800
Met	Gln	Asn	Lys	Val	Gly	Ala	Leu	Asn	Glu	His	Leu	Val	Thr	Gly	Asn
				805					810					815	
Ala	Asp	Lys	Asn	Lys	Ser	Val	His	Tyr	Gln	Ala	Leu	Thr	Ala	Asp	Arg
			820					825					830		
Glu	Trp	Val	Arg	Ser	Arg	Thr	Gly	Leu	Gly	Val	Asn	Pro	Tyr	Asp	Thr
	835						840					845			
Lys	Ser	Val	Lys	Phe	Ala	Gln	Gln	Val	Ala	Leu	Glu	Ala	Lys	Thr	Val
	850					855					860				
Ala	Asp	Ile	Ala	Asn	Ala	Leu	Ala	Ser	Ala	Tyr	Pro	Glu	Leu	Lys	Val
865					870					875					880
Ser	His	Ile	Lys	Val	Thr	Pro	Leu	Asp	Ser	Arg	Leu	Asn	Ala	Pro	Ala
				885					890					895	
Ala	Glu	Val	Val	Lys	Ala	Phe	Arg	Gln	Gly	Asn	Arg	Asp	Val	Ala	Ser
			900					905					910		
Ser	Gln	Pro	Lys	Ala	Asp	Ser	Val	Asn	Gln	Val	Lys	Glu	Thr	Pro	Val
		915					920					925			
Thr	Lys	Gln	Glu	Pro	Val	Thr	Ser	Thr	Val	Gln	Thr	Lys	Thr	Pro	Val
	930					935					940				
Ser	Glu	Ser	Val	Lys	Thr	Glu	Pro	Thr	Thr	Lys	Glu	Ser	Ser	Pro	Gln

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945		950		955		960									
Ala	Ile	Lys	Glu	Pro	Val	Asn	Gln	Ser	Glu	Lys	Gln	Asp	Val	Asn	Leu
			965						970					975	
Thr	Asn	Glu	Asp	Asn	Ile	Lys	Gln	Pro	Thr	Glu	Ser	Val	Lys	Glu	Thr
			980					985					990		
Glu	Thr	Ser	Thr	Lys	Glu	Ser	Thr	Val	Thr	Glu	Glu	Leu	Lys	Glu	Gly
		995					1000					1005			
Ile	Asp	Ala	Val	Tyr	Pro	Ser	Leu	Val	Gly	Thr	Ala	Asp	Ser	Lys	Ala
	1010					1015					1020				
Glu	Gly	Ile	Lys	Asn	Tyr	Phe	Lys	Leu	Ser	Phe	Thr	Leu	Pro	Glu	Glu
1025				1030						1035					1040
Gln	Lys	Ser	Arg	Thr	Val	Gly	Ser	Glu	Ala	Pro	Leu	Lys	Asp	Val	Ala
			1045						1050					1055	
Gln	Ala	Leu	Ser	Ser	Arg	Ala	Arg	Tyr	Glu	Leu	Phe	Thr	Glu	Lys	Glu
		1060						1065					1070		
Thr	Ala	Asn	Pro	Ala	Phe	Asn	Gly	Glu	Val	Ile	Lys	Arg	Tyr	Lys	Glu
	1075						1080					1085			
Leu	Met	Glu	His	Gly	Glu	Gly	Ile	Ala	Asp	Ile	Leu	Arg	Ser	Arg	Leu
	1090					1095					1100				
Ala	Lys	Phe	Leu	Asn	Thr	Lys	Asp	Val	Gly	Lys	Arg	Phe	Ala	Gln	Gly
1105				1110						1115					1120
Thr	Glu	Ala	Asn	Arg	Trp	Val	Gly	Gly	Lys	Leu	Leu	Asn	Ile	Val	Glu
			1125					1130						1135	
Gln	Asp	Gly	Asp	Thr	Phe	Lys	Tyr	Asn	Glu	Gln	Leu	Leu	Gln	Thr	Ala
		1140						1145					1150		
Val	Leu	Ala	Gly	Leu	Gln	Trp	Arg	Leu	Thr	Ala	Thr	Ser	Asn	Thr	Ala
	1155						1160					1165			
Ile	Lys	Asp	Ala	Lys	Asp	Val	Ala	Ala	Ile	Thr	Gly	Ile	Asp	Gln	Ala
	1170				1175						1180				
Leu	Leu	Pro	Glu	Gly	Leu	Val	Glu	Gln	Phe	Asp	Thr	Gly	Met	Thr	Leu
1185				1190					1195						1200
Thr	Glu	Ala	Val	Ser	Ser	Leu	Ala	Gln	Lys	Ile	Glu	Ser	Tyr	Trp	Gly

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1205		1210		1215
Leu Ser Arg Asn Pro Asn Ala Pro Leu Gly Tyr Thr Lys Gly Ile Pro	1220	1225	1230	
Thr Ala Met Ala Ala Glu Ile Leu Ala Ala Phe Val Glu Ser Thr Asp	1235	1240	1245	
Val Val Glu Asn Ile Val Asp Met Ser Glu Ile Asp Pro Asp Asn Lys	1250	1255	1260	
Lys Thr Ile Gly Leu Tyr Thr Ile Thr Glu Leu Asp Ser Phe Asp Pro	1265	1270	1275	1280
Ile Asn Ser Phe Pro Thr Ala Ile Glu Glu Ala Val Leu Val Asn Pro	1285	1290	1295	
Thr Glu Lys Met Phe Phe Gly Asp Asp Ile Pro Pro Val Ala Asn Thr	1300	1305	1310	
Gln Leu Arg Asn Pro Ala Val Arg Asn Thr Pro Glu Gln Lys Ala Ala	1315	1320	1325	
Leu Lys Ala Glu Gln Ala Thr Glu Phe Tyr Val His Thr Pro Met Val	1330	1335	1340	
Gln Phe Tyr Glu Thr Leu Gly Lys Asp Arg Ile Leu Glu Leu Met Gly	1345	1350	1355	1360
Ala Gly Thr Leu Asn Lys Glu Leu Leu Asn Asp Asn His Ala Lys Ser	1365	1370	1375	
Leu Glu Gly Lys Asn Arg Ser Val Glu Asp Ser Tyr Asn Gln Leu Phe	1380	1385	1390	
Ser Val Ile Glu Gln Val Arg Ala Gln Ser Glu Asp Ile Ser Thr Val	1395	1400	1405	
Pro Ile His Tyr Ala Tyr Asn Met Thr Arg Val Gly Arg Met Gln Met	1410	1415	1420	
Leu Gly Lys Tyr Asn Pro Gln Ser Ala Lys Leu Val Arg Glu Ala Ile	1425	1430	1435	1440
Leu Pro Thr Lys Ala Thr Leu Asp Leu Ser Asn Gln Asn Asn Glu Asp	1445	1450	1455	
Phe Ser Ala Phe Gln Leu Gly Leu Ala Gln Ala Leu Asp Ile Lys Val				

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1460	1465	1470
His Thr Met Thr Arg Glu Val Met Ser Asp Glu Leu Thr Lys Leu Leu		
1475	1480	1485
Glu Gly Asn Leu Lys Pro Ala Ile Asp Met Met Val Glu Phe Asn Thr		
1490	1495	1500
Thr Gly Ser Leu Pro Glu Asn Ala Val Asp Val Leu Asn Thr Ala Leu		
1505	1510	1515
Gly Asp Arg Lys Ser Phe Val Ala Leu Met Ala Leu Met Glu Tyr Ser		
1525	1530	1535
Arg Tyr Leu Val Ala Glu Asp Lys Ser Ala Phe Val Thr Pro Leu Tyr		
1540	1545	1550
Val Glu Ala Asp Gly Val Thr Asn Gly Pro Ile Asn Ala Met Met Leu		
1555	1560	1565
Met Thr Gly Gly Leu Phe Thr Pro Asp Trp Ile Arg Asn Ile Ala Lys		
1570	1575	1580
Gly Gly Leu Phe Ile Gly Ser Pro Asn Lys Thr Met Asn Glu His Arg		
1585	1590	1595
Ser Thr Ala Asp Asn Asn Asp Leu Tyr Gln Ala Ser Thr Asn Ala Leu		
1605	1610	1615
Met Glu Ser Leu Gly Lys Leu Arg Ser Asn Tyr Ala Ser Asn Met Pro		
1620	1625	1630
Ile Gln Ser Gln Ile Asp Ser Leu Leu Ser Leu Met Asp Leu Phe Leu		
1635	1640	1645
Pro Asp Ile Asn Leu Gly Glu Asn Gly Ala Leu Glu Leu Lys Arg Gly		
1650	1655	1660
Ile Ala Lys Asn Pro Leu Thr Ile Thr Ile Tyr Gly Ser Gly Ala Arg		
1665	1670	1675
Gly Ile Ala Gly Lys Leu Val Ser Ser Val Thr Asp Ala Ile Tyr Glu		
1685	1690	1695
Arg Met Ser Asp Val Leu Lys Ala Arg Ala Lys Asp Pro Asn Ile Ser		
1700	1705	1710
Ala Ala Met Ala Met Phe Gly Lys Gln Ala Ala Ser Glu Ala His Ala		

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1715	1720	1725
Glu Glu Leu Leu Ala Arg Phe Leu Lys Asp Met Glu Thr Leu Thr Ser		
1730	1735	1740
Thr Val Pro Val Lys Arg Lys Gly Val Leu Glu Leu Gln Ser Thr Gly		
1745	1750	1755 1760
Thr Gly Ala Lys Gly Lys Ile Asn Pro Lys Thr Tyr Thr Ile Lys Gly		
1765	1770	1775
Glu Gln Leu Lys Ala Leu Gln Glu Asn Met Leu His Phe Phe Val Glu		
1780	1785	1790
Pro Leu Arg Asn Gly Ile Thr Gln Thr Val Gly Glu Ser Leu Val Tyr		
1795	1800	1805
Ser Thr Glu Gln Leu Gln Lys Ala Thr Gln Ile Gln Ser Val Val Leu		
1810	1815	1820
Glu Asp Met Phe Lys Gln Arg Val Gln Glu Lys Leu Ala Glu Lys Ala		
1825	1830	1835 1840
Lys Asp Pro Thr Trp Lys Lys Gly Asp Phe Leu Thr Gln Lys Glu Leu		
1845	1850	1855
Asn Asp Ile Gln Ala Ser Leu Asn Asn Leu Ala Pro Met Ile Glu Thr		
1860	1865	1870
Gly Ser Gln Thr Phe Tyr Ile Ala Gly Ser Glu Asn Ala Glu Val Ala		
1875	1880	1885
Asn Gln Val Leu Ala Thr Asn Leu Asp Asp Arg Met Arg Val Pro Met		
1890	1895	1900
Ser Ile Tyr Ala Pro Ala Gln Ala Gly Val Ala Gly Ile Pro Phe Met		
1905	1910	1915 1920
Thr Ile Gly Thr Gly Asp Gly Met Met Met Gln Thr Leu Ser Thr Met		
1925	1930	1935
Lys Gly Ala Pro Lys Asn Thr Leu Lys Ile Phe Asp Gly Met Asn Ile		
1940	1945	1950
Gly Leu Asn Asp Ile Thr Asp Ala Ser Arg Lys Ala Asn Glu Ala Val		
1955	1960	1965
Tyr Thr Ser Trp Gln Gly Asn Pro Ile Lys Asn Val Tyr Glu Ser Tyr		

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1970	1975	1980
Ala Lys Phe Met Lys Asn Val Asp Phe Ser Lys Leu Ser Pro Glu Ala 1985	1990	1995 2000
Leu Glu Ala Ile Gly Lys Ser Ala Leu Glu Tyr Asp Gln Arg Glu Asn 2005	2010	2015
Ala Thr Val Asp Asp Ile Ala Asn Ala Ala Ser Leu Ile Glu Arg Asn 2020	2025	2030
Leu Arg Asn Ile Ala Leu Gly Val Asp Ile Arg His Lys Val Leu Asp 2035	2040	2045
Lys Val Asn Leu Ser Ile Asp Gln Met Ala Ala Val Gly Ala Pro Tyr 2050	2055	2060
Gln Asn Asn Gly Lys Ile Asp Leu Ser Asn Met Thr Pro Glu Gln Gln 2065	2070	2075 2080
Ala Asp Glu Leu Asn Lys Leu Phe Arg Glu Glu Leu Glu Ala Arg Lys 2085	2090	2095
Gln Lys Val Ala Lys Ala Arg Ala Glu Val Lys Glu Glu Thr Val Ser 2100	2105	2110
Glu Lys Glu Pro Val Asn Pro Asp Phe Gly Met Val Gly Arg Glu His 2115	2120	2125
Lys Ala Ser Gly Val Arg Ile Leu Ser Ala Thr Ala Ile Arg Asn Leu 2130	2135	2140
Ala Lys Ile Ser Asn Leu Pro Ser Thr Gln Ala Ala Thr Leu Ala Glu 2145	2150	2155 2160
Ile Gln Lys Ser Leu Ala Ala Lys Asp Tyr Lys Ile Ile Tyr Gly Thr 2165	2170	2175
Pro Thr Gln Val Ala Glu Tyr Ala Arg Gln Lys Asn Val Thr Glu Leu 2180	2185	2190
Thr Ser Gln Glu Met Glu Glu Ala Gln Ala Gly Asn Ile Tyr Gly Trp 2195	2200	2205
Thr Asn Phe Asp Asp Lys Thr Ile Tyr Leu Val Ser Pro Ser Met Glu 2210	2215	2220
Thr Leu Ile His Glu Leu Val His Ala Ser Thr Phe Glu Glu Val Tyr		

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2225	2230	2235	2240
Ser Phe Tyr Gln Gly Asn Glu Val Ser Pro Thr Ser Lys Gln Ala Ile	2245	2250	2255
Glu Asn Leu Glu Gly Leu Met Glu Gln Phe Arg Ser Leu Asp Ile Ser	2260	2265	2270
Lys Asp Ser Pro Glu Met Arg Glu Ala Tyr Ala Asp Ala Ile Ala Thr	2275	2280	2285
Ile Glu Gly His Leu Ser Asn Gly Phe Val Asp Pro Ala Ile Ser Lys	2290	2295	2300
Ala Ala Ala Leu Asn Glu Phe Met Ala Trp Gly Leu Ala Asn Arg Ala	2305	2310	2315
Leu Ala Ala Lys Gln Lys Arg Thr Ser Ser Leu Val Gln Met Val Lys	2325	2330	2335
Asp Val Tyr Gln Ala Ile Lys Lys Leu Ile Trp Gly Arg Lys Gln Ala	2340	2345	2350
Pro Ala Leu Gly Glu Asp Met Phe Ser Asn Leu Leu Phe Asn Ser Ala	2355	2360	2365
Ile Leu Met Arg Ser Gln Pro Thr Thr Gln Ala Val Ala Lys Asp Gly	2370	2375	2380
Thr Leu Phe His Ser Lys Ala Tyr Gly Asn Asn Glu Arg Leu Ser Gln	2385	2390	2395
Leu Asn Gln Thr Phe Asp Lys Leu Val Thr Asp Tyr Leu Arg Thr Asp	2405	2410	2415
Pro Val Thr Glu Val Glu Arg Arg Gly Asn Val Ala Asn Ala Leu Met	2420	2425	2430
Ser Ala Thr Arg Leu Val Arg Asp Val Gln Ser His Gly Phe Asn Met	2435	2440	2445
Thr Ala Gln Glu Gln Ser Val Phe Gln Met Val Thr Ala Ala Leu Ala	2450	2455	2460
Thr Glu Ala Ala Ile Asp Pro His Ala Met Ala Arg Ala Gln Glu Leu	2465	2470	2475
Tyr Thr His Val Met Lys His Leu Thr Val Glu His Phe Met Ala Asp			

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2485 2490 2495

Pro Asp Ser Thr Asn Pro Ala Asp Arg Tyr Tyr Ala Gln Gln Lys Tyr
2500 2505 2510

Asp Thr Ile Ser Gly Ala Asn Leu Val Glu Val Asp Ala Lys Gly Arg
2515 2520 2525

Thr Ser Leu Leu Pro Thr Phe Leu Gly Leu Ala Met Val Asn Glu Glu
2530 2535 2540

Leu Arg Ser Ile Ile Lys Glu Met Pro Val Pro Lys Ala Asp Lys Lys
2545 2550 2555 2560

Leu Gly Asn Asp Ile Asp Thr Leu Leu Thr Asn Ala Gly Thr Gln Val
2565 2570 2575

Met Glu Ser Leu Asn Arg Arg Met Ala Gly Asp Gln Lys Ala Thr Asn
2580 2585 2590

Val Gln Asp Ser Ile Asp Ala Leu Ser Glu Thr Ile Met Ala Ala Ala
2595 2600 2605

Leu Lys Arg Glu Ser Phe Tyr Asp Ala Val Ala Thr Pro Thr Gly Asn
2610 2615 2620

Phe Ile Asp Arg Ala Asn Gln Tyr Val Thr Asp Ser Ile Glu Arg Leu
2625 2630 2635 2640

Ser Glu Thr Val Ile Glu Lys Ala Asp Lys Val Ile Ala Asn Pro Ser
2645 2650 2655

Asn Ile Ala Ala Lys Gly Val Ala His Leu Ala Lys Leu Thr Ala Ala
2660 2665 2670

Ile Ala Ser Glu Lys Gln Gly Glu Ile Val Ala Gln Gly Val Met Thr
2675 2680 2685

Ala Met Asn Gln Gly Lys Val Trp Gln Pro Phe His Asp Leu Val Asn
2690 2695 2700

Asp Ile Val Gly Arg Thr Lys Thr Asn Ala Asn Val Tyr Asp Leu Ile
2705 2710 2715 2720

Lys Leu Val Lys Ser Gln Ile Ser Gln Asp Arg Gln Gln Phe Arg Glu
2725 2730 2735

His Leu Pro Thr Val Ile Ala Gly Lys Phe Ser Arg Lys Leu Thr Asp

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2740		2745		2750
Thr Glu Trp Ser Ala Met His Thr Gly Leu Gly Lys Thr Asp Leu Ala	2755	2760	2765	
Val Leu Arg Glu Thr Met Ser Met Ala Glu Ile Arg Asp Leu Leu Ser	2770	2775	2780	
Ser Ser Lys Lys Val Lys Asp Glu Ile Ser Thr Leu Glu Lys Glu Ile	2785	2790	2795	2800
Gln Asn Gln Ala Gly Arg Asn Trp Asn Leu Val Gln Lys Lys Ser Lys	2805	2810	2815	
Gln Leu Ala Gln Tyr Met Ile Met Gly Glu Val Gly Asn Asn Leu Leu	2820	2825	2830	
Arg Asn Ala His Ala Ile Ser Arg Leu Leu Gly Glu Arg Ile Thr Asn	2835	2840	2845	
Gly Pro Val Ala Asp Val Ala Ala Ile Asp Lys Leu Ile Thr Leu Tyr	2850	2855	2860	
Ser Leu Glu Leu Met Asn Lys Ser Asp Arg Asp Leu Leu Ser Glu Leu	2865	2870	2875	2880
Ala Gln Ser Glu Val Glu Gly Met Glu Phe Ser Ile Ala Tyr Met Val	2885	2890	2895	
Gly Gln Arg Thr Glu Glu Met Arg Lys Ala Lys Gly Asp Asn Arg Thr	2900	2905	2910	
Leu Leu Asn His Phe Lys Gly Tyr Ile Pro Val Glu Asn Gln Gln Gly	2915	2920	2925	
Val Asn Leu Ile Ile Ala Asp Asp Lys Glu Phe Ala Lys Leu Asn Ser	2930	2935	2940	
Gln Ser Phe Thr Arg Ile Gly Thr Tyr Gln Gly Ser Thr Gly Phe Arg	2945	2950	2955	2960
Thr Gly Ser Lys Gly Tyr Tyr Phe Ser Pro Val Ala Ala Arg Ala Pro	2965	2970	2975	
Tyr Ser Gln Gly Ile Leu Gln Asn Val Arg Asn Thr Ala Gly Gly Val	2980	2985	2990	
Asp Ile Gly Thr Gly Phe Thr Leu Gly Thr Met Val Ala Gly Arg Ile				

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2995	3000	3005															
Thr	Asp	Lys	Pro	Thr	Val	Glu	Arg	Ile	Thr	Lys	Ala	Leu	Ala	Lys	Gly		
3010					3015					3020							
Glu	Arg	Gly	Arg	Glu	Pro	Leu	Met	Pro	Ile	Tyr	Asn	Ser	Lys	Gly	Gln		
3025				3030					3035						3040		
Val	Val	Ala	Tyr	Glu	Gln	Ser	Val	Asp	Pro	Asn	Met	Leu	Lys	His	Leu		
			3045					3050						3055			
Asn	Gln	Asp	Asn	His	Phe	Ala	Lys	Met	Val	Gly	Val	Trp	Arg	Gly	Arg		
		3060						3065					3070				
Gln	Val	Glu	Glu	Ala	Lys	Ala	Gln	Arg	Phe	Asn	Asp	Ile	Leu	Ile	Glu		
	3075						3080					3085					
Gln	Leu	His	Ala	Met	Tyr	Glu	Lys	Asp	Ile	Lys	Asp	Ser	Ser	Ala	Asn		
3090					3095						3100						
Lys	Ser	Gln	Tyr	Val	Asn	Leu	Leu	Gly	Lys	Ile	Asp	Asp	Pro	Val	Leu		
3105				3110						3115					3120		
Ala	Asp	Ala	Ile	Asn	Leu	Met	Asn	Ile	Glu	Thr	Arg	His	Lys	Ala	Glu		
			3125						3130					3135			
Glu	Leu	Phe	Gly	Lys	Asp	Glu	Leu	Trp	Val	Arg	Arg	Asp	Met	Leu	Asn		
		3140						3145					3150				
Asp	Ala	Leu	Gly	Tyr	Arg	Ala	Ala	Ser	Ile	Gly	Asp	Val	Trp	Thr	Gly		
	3155						3160					3165					
Asn	Ser	Arg	Trp	Ser	Pro	Ser	Thr	Leu	Asp	Thr	Val	Lys	Lys	Met	Phe		
3170					3175						3180						
Leu	Gly	Ala	Phe	Gly	Asn	Lys	Ala	Tyr	His	Val	Val	Met	Asn	Ala	Glu		
3185				3190						3195					3200		
Asn	Thr	Ile	Gln	Asn	Leu	Val	Lys	Asp	Ala	Lys	Thr	Val	Ile	Val	Val		
			3205						3210					3215			
Lys	Ser	Val	Val	Val	Pro	Ala	Val	Asn	Phe	Leu	Ala	Asn	Ile	Tyr	Gln		
		3220						3225					3230				
Met	Ile	Gly	Arg	Gly	Val	Pro	Val	Lys	Asp	Ile	Ala	Val	Asn	Ile	Pro		
	3235						3240					3245					
Arg	Lys	Thr	Ser	Glu	Ile	Asn	Gln	Tyr	Ile	Lys	Ser	Arg	Leu	Arg	Gln		

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3250	3255	3260
Ile Asp Ala Glu Ala Glu Leu Arg Ala Ala Glu Gly Asn Pro Asn Leu		
3265	3270	3275 3280
Val Arg Lys Leu Lys Thr Glu Ile Gln Ser Ile Thr Asp Ser His Arg		
	3285	3290 3295
Arg Met Ser Ile Trp Pro Leu Ile Glu Ala Gly Glu Phe Ser Ser Ile		
	3300	3305 3310
Ala Asp Ala Gly Ile Ser Arg Asp Asp Leu Leu Val Ala Glu Gly Lys		
	3315	3320 3325
Ile His Glu Tyr Met Glu Lys Leu Ala Asn Lys Leu Pro Glu Lys Val		
	3330	3335 3340
Arg Asn Ala Gly Arg Tyr Ala Leu Ile Ala Lys Asp Thr Ala Leu Phe		
	3345	3350 3355 3360
Gln Gly Ile Gln Lys Thr Val Glu Tyr Ser Asp Phe Ile Ala Lys Ala		
	3365	3370 3375
Ile Ile Tyr Asp Asp Leu Val Lys Arg Lys Lys Lys Ser Ser Ser Glu		
	3380	3385 3390
Ala Leu Gly Gln Val Thr Glu Glu Phe Ile Asn Tyr Asp Arg Leu Pro		
	3395	3400 3405
Gly Arg Phe Arg Gly Tyr Met Glu Ser Met Gly Leu Met Trp Phe Tyr		
	3410	3415 3420
Asn Phe Lys Ile Arg Ser Ile Lys Val Ala Met Ser Met Ile Arg Asn		
	3425	3430 3435 3440
Asn Pro Val His Ser Leu Ile Ala Thr Val Val Pro Ala Pro Thr Met		
	3445	3450 3455
Phe Gly Asn Val Gly Leu Pro Ile Gln Asp Asn Met Leu Thr Met Leu		
	3460	3465 3470
Ala Glu Gly Arg Leu Asp Tyr Ser Leu Gly Phe Gly Gln Gly Leu Arg		
	3475	3480 3485
Ala Pro Thr Leu Asn Pro Trp Phe Asn Leu Thr His		
	3490	3495 3500

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<210> 3

<211> 3318

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
Primer

<400> 3

gaaagtacag ttacagaaga attaaaagaa ggtattgatg ctgtttaccc ttcattggta 60
ggtactgctg attctaaagc agaggggtatt aagaactatt tcaaattgtc ctttacctta 120
ccagaagaac agaaatcccc tactgttggt tcagaagcac ctctaaaaga tgtagcccaa 180
gctctgtctt ctctgtctcg ttatgaactc tttactgaga aagaaactgc taaccctgct 240
tttaatgggg aagttattaa gcgatacaaa gaactcatgg aacatgggga aggtattgct 300
gatattcttc gctcccgctt ggctaagttc cttaacacta aggatggttg taaacgtttt 360
gctcaaggta cagaagccaa ccgttgggta ggtggtaagt tacttaacat tgttgagcag 420
gatggggata cctttaagta caacgaacaa ttgctacaga ctgctgtatt agcaggtctt 480
caatggagac ttactgctac cagcaatact gctatcaaag atgcaaaaaga tgttgctgct 540
attactggta ttgaccaagc tctgctgcca gaaggtttag tagagcaatt tgatactggt 600
atgacactca ctgaagcagt tagttccctg gctcagaaaa ttgagtctta ctggggatta 660
tctcgtaatc caaatgctcc attgggctat accaaaggca tccctacagc aatggctgct 720
gaaattctgg ctgcatttgt agagtctact gatggtgtag agaacatcgt ggatatgtca 780
gaaattgacc cagataacaa gaagactatt ggtctgtaca ccattactga actggattcc 840
ttcgacccaa ttaatagctt ccctactgct attgaagaag ctgttttagt gaatcctaca 900
gagaagatgt tctttgggtga tgacattcct cctgtagcta atactcagct tcgtaaccct 960
gctgttcgta atactccaga acagaaggct gcattgaaag cagagcaggc tacagagttc 1020
tatgtacaca cccaatgggt tcaattctat gagacgtag gtaaagaccg tattctcgaa 1080
ctgatgggtg ctggtactct gaataaagag ttacttaatg ataaccatgc taaatctctg 1140

Epicentre-00005 seqlist (Nov).txt

gaaggtaaga accgttcagt agaggactct tacaaccaac tggtctccgt cattgagcag 1200
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Epicentre-00005 seqlist (Nov).txt

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 aaagtcgcta aggctagg 3318

<210> 4

<211> 1107

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 Peptide

<400> 4

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 1 5 10 15

Tyr Pro Ser Leu Val Gly Thr Ala Asp Ser Lys Ala Glu Gly Ile Lys
 20 25 30

Asn Tyr Phe Lys Leu Ser Phe Thr Leu Pro Glu Glu Gln Lys Ser Arg
 35 40 45

Thr Val Gly Ser Glu Ala Pro Leu Lys Asp Val Ala Gln Ala Leu Ser

Epicentre-00005 seqlist (Nov).txt

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Ala	Phe	Asn	Gly	Glu	Val	Ile	Lys	Arg	Tyr	Lys	Glu	Leu	Met	Glu	His	
				85					90					95		
Gly	Glu	Gly	Ile	Ala	Asp	Ile	Leu	Arg	Ser	Arg	Leu	Ala	Lys	Phe	Leu	
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Asn	Thr	Lys	Asp	Val	Gly	Lys	Arg	Phe	Ala	Gln	Gly	Thr	Glu	Ala	Asn	
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Arg	Trp	Val	Gly	Gly	Lys	Leu	Leu	Asn	Ile	Val	Glu	Gln	Asp	Gly	Asp	
	130					135					140					
Thr	Phe	Lys	Tyr	Asn	Glu	Gln	Leu	Leu	Gln	Thr	Ala	Val	Leu	Ala	Gly	
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Leu	Gln	Trp	Arg	Leu	Thr	Ala	Thr	Ser	Asn	Thr	Ala	Ile	Lys	Asp	Ala	
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Lys	Asp	Val	Ala	Ala	Ile	Thr	Gly	Ile	Asp	Gln	Ala	Leu	Leu	Pro	Glu	
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Gly	Leu	Val	Glu	Gln	Phe	Asp	Thr	Gly	Met	Thr	Leu	Thr	Glu	Ala	Val	
		195					200					205				
Ser	Ser	Leu	Ala	Gln	Lys	Ile	Glu	Ser	Tyr	Trp	Gly	Leu	Ser	Arg	Asn	
	210					215					220					
Pro	Asn	Ala	Pro	Leu	Gly	Tyr	Thr	Lys	Gly	Ile	Pro	Thr	Ala	Met	Ala	
225					230					235					240	
Ala	Glu	Ile	Leu	Ala	Ala	Phe	Val	Glu	Ser	Thr	Asp	Val	Val	Glu	Asn	
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Ile	Val	Asp	Met	Ser	Glu	Ile	Asp	Pro	Asp	Asn	Lys	Lys	Thr	Ile	Gly	
			260					265					270			
Leu	Tyr	Thr	Ile	Thr	Glu	Leu	Asp	Ser	Phe	Asp	Pro	Ile	Asn	Ser	Phe	
		275					280					285				
Pro	Thr	Ala	Ile	Glu	Glu	Ala	Val	Leu	Val	Asn	Pro	Thr	Glu	Lys	Met	
	290					295					300					
Phe	Phe	Gly	Asp	Asp	Ile	Pro	Pro	Val	Ala	Asn	Thr	Gln	Leu	Arg	Asn	

Epicentre-00005 seqlist (Nov).txt

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305          310          315          320
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          325          330          335
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Thr Leu Gly Lys Asp Arg Ile Leu Glu Leu Met Gly Ala Gly Thr Leu
          355          360          365
Asn Lys Glu Leu Leu Asn Asp Asn His Ala Lys Ser Leu Glu Gly Lys
          370          375          380
Asn Arg Ser Val Glu Asp Ser Tyr Asn Gln Leu Phe Ser Val Ile Glu
385          390          395          400
Gln Val Arg Ala Gln Ser Glu Asp Ile Ser Thr Val Pro Ile His Tyr
          405          410          415
Ala Tyr Asn Met Thr Arg Val Gly Arg Met Gln Met Leu Gly Lys Tyr
          420          425          430
Asn Pro Gln Ser Ala Lys Leu Val Arg Glu Ala Ile Leu Pro Thr Lys
          435          440          445
Ala Thr Leu Asp Leu Ser Asn Gln Asn Asn Glu Asp Phe Ser Ala Phe
          450          455          460
Gln Leu Gly Leu Ala Gln Ala Leu Asp Ile Lys Val His Thr Met Thr
465          470          475          480
Arg Glu Val Met Ser Asp Glu Leu Thr Lys Leu Leu Glu Gly Asn Leu
          485          490          495
Lys Pro Ala Ile Asp Met Met Val Glu Phe Asn Thr Thr Gly Ser Leu
          500          505          510
Pro Glu Asn Ala Val Asp Val Leu Asn Thr Ala Leu Gly Asp Arg Lys
          515          520          525
Ser Phe Val Ala Leu Met Ala Leu Met Glu Tyr Ser Arg Tyr Leu Val
          530          535          540
Ala Glu Asp Lys Ser Ala Phe Val Thr Pro Leu Tyr Val Glu Ala Asp
545          550          555          560
Gly Val Thr Asn Gly Pro Ile Asn Ala Met Met Leu Met Thr Gly Gly

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Epicentre-00005 seqlist (Nov).txt
565 570 575

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Ile	Gly	Ser	Pro	Asn	Lys	Thr	Met	Asn	Glu	His	Arg	Ser	Thr	Ala	Asp
		595					600					605			
Asn	Asn	Asp	Leu	Tyr	Gln	Ala	Ser	Thr	Asn	Ala	Leu	Met	Glu	Ser	Leu
	610					615					620				
Gly	Lys	Leu	Arg	Ser	Asn	Tyr	Ala	Ser	Asn	Met	Pro	Ile	Gln	Ser	Gln
625					630					635					640
Ile	Asp	Ser	Leu	Leu	Ser	Leu	Met	Asp	Leu	Phe	Leu	Pro	Asp	Ile	Asn
			645						650					655	
Leu	Gly	Glu	Asn	Gly	Ala	Leu	Glu	Leu	Lys	Arg	Gly	Ile	Ala	Lys	Asn
			660					665					670		
Pro	Leu	Thr	Ile	Thr	Ile	Tyr	Gly	Ser	Gly	Ala	Arg	Gly	Ile	Ala	Gly
		675					680					685			
Lys	Leu	Val	Ser	Ser	Val	Thr	Asp	Ala	Ile	Tyr	Glu	Arg	Met	Ser	Asp
	690					695					700				
Val	Leu	Lys	Ala	Arg	Ala	Lys	Asp	Pro	Asn	Ile	Ser	Ala	Ala	Met	Ala
705					710					715					720
Met	Phe	Gly	Lys	Gln	Ala	Ala	Ser	Glu	Ala	His	Ala	Glu	Glu	Leu	Leu
				725					730					735	
Ala	Arg	Phe	Leu	Lys	Asp	Met	Glu	Thr	Leu	Thr	Ser	Thr	Val	Pro	Val
			740					745					750		
Lys	Arg	Lys	Gly	Val	Leu	Glu	Leu	Gln	Ser	Thr	Gly	Thr	Gly	Ala	Lys
		755					760					765			
Gly	Lys	Ile	Asn	Pro	Lys	Thr	Tyr	Thr	Ile	Lys	Gly	Glu	Gln	Leu	Lys
	770					775					780				
Ala	Leu	Gln	Glu	Asn	Met	Leu	His	Phe	Phe	Val	Glu	Pro	Leu	Arg	Asn
785					790					795					800
Gly	Ile	Thr	Gln	Thr	Val	Gly	Glu	Ser	Leu	Val	Tyr	Ser	Thr	Glu	Gln
				805					810					815	
Leu	Gln	Lys	Ala	Thr	Gln	Ile	Gln	Ser	Val	Val	Leu	Glu	Asp	Met	Phe

Epicentre-00005 seqlist (Nov).txt

820								825							830
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	835						840					845			
Trp	Lys	Lys	Gly	Asp	Phe	Leu	Thr	Gln	Lys	Glu	Leu	Asn	Asp	Ile	Gln
	850					855					860				
Ala	Ser	Leu	Asn	Asn	Leu	Ala	Pro	Met	Ile	Glu	Thr	Gly	Ser	Gln	Thr
865					870					875				880	
Phe	Tyr	Ile	Ala	Gly	Ser	Glu	Asn	Ala	Glu	Val	Ala	Asn	Gln	Val	Leu
				885					890					895	
Ala	Thr	Asn	Leu	Asp	Asp	Arg	Met	Arg	Val	Pro	Met	Ser	Ile	Tyr	Ala
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Lys	Asn	Thr	Leu	Lys	Ile	Phe	Asp	Gly	Met	Asn	Ile	Gly	Leu	Asn	Asp
945					950					955					960
Ile	Thr	Asp	Ala	Ser	Arg	Lys	Ala	Asn	Glu	Ala	Val	Tyr	Thr	Ser	Trp
				965					970					975	
Gln	Gly	Asn	Pro	Ile	Lys	Asn	Val	Tyr	Glu	Ser	Tyr	Ala	Lys	Phe	Met
			980					985					990		
Lys	Asn	Val	Asp	Phe	Ser	Lys	Leu	Ser	Pro	Glu	Ala	Leu	Glu	Ala	Ile
		995					1000					1005			
Gly	Lys	Ser	Ala	Leu	Glu	Tyr	Asp	Gln	Arg	Glu	Asn	Ala	Thr	Val	Asp
	1010					1015					1020				
Asp	Ile	Ala	Asn	Ala	Ala	Ser	Leu	Ile	Glu	Arg	Asn	Leu	Arg	Asn	Ile
1025				1030					1035					1040	
Ala	Leu	Gly	Val	Asp	Ile	Arg	His	Lys	Val	Leu	Asp	Lys	Val	Asn	Leu
			1045					1050						1055	
Ser	Ile	Asp	Gln	Met	Ala	Ala	Val	Gly	Ala	Pro	Tyr	Gln	Asn	Asn	Gly
		1060					1065					1070			
Lys	Ile	Asp	Leu	Ser	Asn	Met	Thr	Pro	Glu	Gln	Gln	Ala	Asp	Glu	Leu

Epicentre-00005 seqlist (Nov).txt

1075

1080

1085

Asn Lys Leu Phe Arg Glu Glu Leu Glu Ala Arg Lys Gln Lys Val Ala
 1090 1095 1100

Lys Ala Arg
 1105

<210> 5

<211> 3432

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 Primer

<400> 5

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 gttacagaag aattaaaaga aggtattgat gctgtttacc cttcattggg aggtactgct 180
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 tctcgtgctc gttatgaact ctttactgag aaagaaactg ctaaccctgc ttttaattggg 360
 gaagttatta agcgatacaa agaactcatg gaacatgggg aaggatttgc tgatattctt 420
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 acagaagcca accgttgggt aggtggtaag ttacttaaca ttgttgagca ggatggggat 540
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Epicentre-00005 seqlist (Nov).txt

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Epicentre-00005 seqlist (Nov).txt

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aaggctaggt aa 3432

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<210> 6

<211> 1143

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic

Epicentre-00005 seqlist (Nov).txt

Peptide

<400> 6

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			20					25					30		
Pro	Ser	Ser	Arg	Ser	Glu	Ser	Thr	Val	Thr	Glu	Glu	Leu	Lys	Glu	Gly
		35					40					45			
Ile	Asp	Ala	Val	Tyr	Pro	Ser	Leu	Val	Gly	Thr	Ala	Asp	Ser	Lys	Ala
	50					55					60				
Glu	Gly	Ile	Lys	Asn	Tyr	Phe	Lys	Leu	Ser	Phe	Thr	Leu	Pro	Glu	Glu
65					70					75					80
Gln	Lys	Ser	Arg	Thr	Val	Gly	Ser	Glu	Ala	Pro	Leu	Lys	Asp	Val	Ala
				85					90					95	
Gln	Ala	Leu	Ser	Ser	Arg	Ala	Arg	Tyr	Glu	Leu	Phe	Thr	Glu	Lys	Glu
			100					105					110		
Thr	Ala	Asn	Pro	Ala	Phe	Asn	Gly	Glu	Val	Ile	Lys	Arg	Tyr	Lys	Glu
		115					120					125			
Leu	Met	Glu	His	Gly	Glu	Gly	Ile	Ala	Asp	Ile	Leu	Arg	Ser	Arg	Leu
	130					135					140				
Ala	Lys	Phe	Leu	Asn	Thr	Lys	Asp	Val	Gly	Lys	Arg	Phe	Ala	Gln	Gly
145					150					155					160
Thr	Glu	Ala	Asn	Arg	Trp	Val	Gly	Gly	Lys	Leu	Leu	Asn	Ile	Val	Glu
				165					170					175	
Gln	Asp	Gly	Asp	Thr	Phe	Lys	Tyr	Asn	Glu	Gln	Leu	Leu	Gln	Thr	Ala
			180					185					190		
Val	Leu	Ala	Gly	Leu	Gln	Trp	Arg	Leu	Thr	Ala	Thr	Ser	Asn	Thr	Ala
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Ile	Lys	Asp	Ala	Lys	Asp	Val	Ala	Ala	Ile	Thr	Gly	Ile	Asp	Gln	Ala
	210					215					220				
Leu	Leu	Pro	Glu	Gly	Leu	Val	Glu	Gln	Phe	Asp	Thr	Gly	Met	Thr	Leu
225					230					235					240

Epicentre-00005 seqlist (Nov).txt

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Thr	Ala	Met	Ala	Ala	Glu	Ile	Leu	Ala	Ala	Phe	Val	Glu	Ser	Thr	Asp	275	280	285
Val	Val	Glu	Asn	Ile	Val	Asp	Met	Ser	Glu	Ile	Asp	Pro	Asp	Asn	Lys	290	295	300
Lys	Thr	Ile	Gly	Leu	Tyr	Thr	Ile	Thr	Glu	Leu	Asp	Ser	Phe	Asp	Pro	305	310	315
Ile	Asn	Ser	Phe	Pro	Thr	Ala	Ile	Glu	Glu	Ala	Val	Leu	Val	Asn	Pro	325	330	335
Thr	Glu	Lys	Met	Phe	Phe	Gly	Asp	Asp	Ile	Pro	Pro	Val	Ala	Asn	Thr	340	345	350
Gln	Leu	Arg	Asn	Pro	Ala	Val	Arg	Asn	Thr	Pro	Glu	Gln	Lys	Ala	Ala	355	360	365
Leu	Lys	Ala	Glu	Gln	Ala	Thr	Glu	Phe	Tyr	Val	His	Thr	Pro	Met	Val	370	375	380
Gln	Phe	Tyr	Glu	Thr	Leu	Gly	Lys	Asp	Arg	Ile	Leu	Glu	Leu	Met	Gly	385	390	395
Ala	Gly	Thr	Leu	Asn	Lys	Glu	Leu	Leu	Asn	Asp	Asn	His	Ala	Lys	Ser	405	410	415
Leu	Glu	Gly	Lys	Asn	Arg	Ser	Val	Glu	Asp	Ser	Tyr	Asn	Gln	Leu	Phe	420	425	430
Ser	Val	Ile	Glu	Gln	Val	Arg	Ala	Gln	Ser	Glu	Asp	Ile	Ser	Thr	Val	435	440	445
Pro	Ile	His	Tyr	Ala	Tyr	Asn	Met	Thr	Arg	Val	Gly	Arg	Met	Gln	Met	450	455	460
Leu	Gly	Lys	Tyr	Asn	Pro	Gln	Ser	Ala	Lys	Leu	Val	Arg	Glu	Ala	Ile	465	470	475
Leu	Pro	Thr	Lys	Ala	Thr	Leu	Asp	Leu	Ser	Asn	Gln	Asn	Asn	Glu	Asp	485	490	495

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Phe	Ser	Ala	Phe	Gln	Leu	Gly	Leu	Ala	Gln	Ala	Leu	Asp	Ile	Lys	Val	500	505	510
His	Thr	Met	Thr	Arg	Glu	Val	Met	Ser	Asp	Glu	Leu	Thr	Lys	Leu	Leu	515	520	525
Glu	Gly	Asn	Leu	Lys	Pro	Ala	Ile	Asp	Met	Met	Val	Glu	Phe	Asn	Thr	530	535	540
Thr	Gly	Ser	Leu	Pro	Glu	Asn	Ala	Val	Asp	Val	Leu	Asn	Thr	Ala	Leu	545	550	555
Gly	Asp	Arg	Lys	Ser	Phe	Val	Ala	Leu	Met	Ala	Leu	Met	Glu	Tyr	Ser	565	570	575
Arg	Tyr	Leu	Val	Ala	Glu	Asp	Lys	Ser	Ala	Phe	Val	Thr	Pro	Leu	Tyr	580	585	590
Val	Glu	Ala	Asp	Gly	Val	Thr	Asn	Gly	Pro	Ile	Asn	Ala	Met	Met	Leu	595	600	605
Met	Thr	Gly	Gly	Leu	Phe	Thr	Pro	Asp	Trp	Ile	Arg	Asn	Ile	Ala	Lys	610	615	620
Gly	Gly	Leu	Phe	Ile	Gly	Ser	Pro	Asn	Lys	Thr	Met	Asn	Glu	His	Arg	625	630	635
Ser	Thr	Ala	Asp	Asn	Asn	Asp	Leu	Tyr	Gln	Ala	Ser	Thr	Asn	Ala	Leu	645	650	655
Met	Glu	Ser	Leu	Gly	Lys	Leu	Arg	Ser	Asn	Tyr	Ala	Ser	Asn	Met	Pro	660	665	670
Ile	Gln	Ser	Gln	Ile	Asp	Ser	Leu	Leu	Ser	Leu	Met	Asp	Leu	Phe	Leu	675	680	685
Pro	Asp	Ile	Asn	Leu	Gly	Glu	Asn	Gly	Ala	Leu	Glu	Leu	Lys	Arg	Gly	690	695	700
Ile	Ala	Lys	Asn	Pro	Leu	Thr	Ile	Thr	Ile	Tyr	Gly	Ser	Gly	Ala	Arg	705	710	715
Gly	Ile	Ala	Gly	Lys	Leu	Val	Ser	Ser	Val	Thr	Asp	Ala	Ile	Tyr	Glu	725	730	735
Arg	Met	Ser	Asp	Val	Leu	Lys	Ala	Arg	Ala	Lys	Asp	Pro	Asn	Ile	Ser	740	745	750

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Ala	Ala	Met	Ala	Met	Phe	Gly	Lys	Gln	Ala	Ala	Ser	Glu	Ala	His	Ala		
		755					760					765					
Glu	Glu	Leu	Leu	Ala	Arg	Phe	Leu	Lys	Asp	Met	Glu	Thr	Leu	Thr	Ser		
	770					775					780						
Thr	Val	Pro	Val	Lys	Arg	Lys	Gly	Val	Leu	Glu	Leu	Gln	Ser	Thr	Gly		
785					790					795					800		
Thr	Gly	Ala	Lys	Gly	Lys	Ile	Asn	Pro	Lys	Thr	Tyr	Thr	Ile	Lys	Gly		
				805					810					815			
Glu	Gln	Leu	Lys	Ala	Leu	Gln	Glu	Asn	Met	Leu	His	Phe	Phe	Val	Glu		
			820					825					830				
Pro	Leu	Arg	Asn	Gly	Ile	Thr	Gln	Thr	Val	Gly	Glu	Ser	Leu	Val	Tyr		
		835					840					845					
Ser	Thr	Glu	Gln	Leu	Gln	Lys	Ala	Thr	Gln	Ile	Gln	Ser	Val	Val	Leu		
	850					855					860						
Glu	Asp	Met	Phe	Lys	Gln	Arg	Val	Gln	Glu	Lys	Leu	Ala	Glu	Lys	Ala		
865					870					875					880		
Lys	Asp	Pro	Thr	Trp	Lys	Lys	Gly	Asp	Phe	Leu	Thr	Gln	Lys	Glu	Leu		
				885					890					895			
Asn	Asp	Ile	Gln	Ala	Ser	Leu	Asn	Asn	Leu	Ala	Pro	Met	Ile	Glu	Thr		
			900					905					910				
Gly	Ser	Gln	Thr	Phe	Tyr	Ile	Ala	Gly	Ser	Glu	Asn	Ala	Glu	Val	Ala		
		915					920					925					
Asn	Gln	Val	Leu	Ala	Thr	Asn	Leu	Asp	Asp	Arg	Met	Arg	Val	Pro	Met		
	930					935					940						
Ser	Ile	Tyr	Ala	Pro	Ala	Gln	Ala	Gly	Val	Ala	Gly	Ile	Pro	Phe	Met		
945					950					955					960		
Thr	Ile	Gly	Thr	Gly	Asp	Gly	Met	Met	Met	Gln	Thr	Leu	Ser	Thr	Met		
				965					970					975			
Lys	Gly	Ala	Pro	Lys	Asn	Thr	Leu	Lys	Ile	Phe	Asp	Gly	Met	Asn	Ile		
			980					985					990				
Gly	Leu	Asn	Asp	Ile	Thr	Asp	Ala	Ser	Arg	Lys	Ala	Asn	Glu	Ala	Val		
		995					1000					1005					

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Tyr Thr Ser Trp Gln Gly Asn Pro Ile Lys Asn Val Tyr Glu Ser Tyr
1010 1015 1020

Ala Lys Phe Met Lys Asn Val Asp Phe Ser Lys Leu Ser Pro Glu Ala
1025 1030 1035 1040

Leu Glu Ala Ile Gly Lys Ser Ala Leu Glu Tyr Asp Gln Arg Glu Asn
1045 1050 1055

Ala Thr Val Asp Asp Ile Ala Asn Ala Ala Ser Leu Ile Glu Arg Asn
1060 1065 1070

Leu Arg Asn Ile Ala Leu Gly Val Asp Ile Arg His Lys Val Leu Asp
1075 1080 1085

Lys Val Asn Leu Ser Ile Asp Gln Met Ala Ala Val Gly Ala Pro Tyr
1090 1095 1100

Gln Asn Asn Gly Lys Ile Asp Leu Ser Asn Met Thr Pro Glu Gln Gln
1105 1110 1115 1120

Ala Asp Glu Leu Asn Lys Leu Phe Arg Glu Glu Leu Glu Ala Arg Lys
1125 1130 1135

Gln Lys Val Ala Lys Ala Arg
1140

<210> 7

<211> 3432

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
Primer

<400> 7

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atgggtcggg atctgtacga cgatgacgat aaggatccga gctcgagatc tgaaagtaca 120
gttacagaag aattaaaga aggtattgat gctgtttacc cttcattggg aggtactgct 180
gattctaaag cagagggtat taagaactat ttcaaattgt cctttacctt accagaagaa 240
cagaaatccc gtactgttgg ttcagaagca cctctaaaag atgtagccca agctctgtct 300

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tctcgtgctc gttatgaact ctttactgag aaagaaactg ctaaccctgc ttttaatggg 360
gaagttatta agcgatacaa agaactcatg gaacatgggg aaggtattgc tgatattctt 420
cgctcccgtc tggctaagtt ccttaacact aaggatgttg gtaaacgttt tgctcaaggt 480
acagaagcca accgttgggt aggtggtaag ttacttaaca ttggtgagca ggatggggat 540
acctttaagt acaacgaaca attgctacag actgctgtat tagcaggtct tcaatggaga 600
cttactgcta ccagcaatac tgctatcaaa gatgcaaaag atgttgctgc tattactggg 660
attgaccaag ctctgctgcc agaaggttta gtagagcaat ttgatactgg tatgacactc 720
actgaagcag ttagttccct ggctcagaaa attgagtctt actggggatt atctcgtaat 780
ccaaatgctc cattgggcta taccaaaggc atccctacag caatggctgc tgaaattctg 840
gctgcatttg tagagtctac tgatgttgta gagaacatcg tggatatgtc agaaattgac 900
ccagataaca agaagactat tggctctgtac accattactg aactggattc cttcgaccca 960
attaatagct tccctactgc tattgaagaa gctgttttag tgaatcctac agagaagatg 1020
ttctttgggtg atgacattcc tcctgtagct aatactcagc ttcgtaaccc tgctgttcgt 1080
aatactccag aacagaaggc tgcattgaaa gcagagcagg ctacagagtt ctatgtacac 1140
acccaatgg ttcaattcta tgagacgtta ggtaaagacc gtattctcga actgatgggt 1200
gctggtactc tgaataaaga gttacttaat gataaccatg ctaaactctt ggaaggtaag 1260
aaccgttcag tagaggactc ttacaaccaa ctgttctccg tcattgagca ggtaagagca 1320
cagagcgaag acatctctac tgtacctatt cactatgcat acaatatgac ccgtgttggt 1380
cgtatgcaga tgtaggtaa atacaatcct caatcagcca aactggttcg tgaggccatc 1440
ttacctacta aagctacttt ggatttatcg aaccagaaca atgaagactt ctctgcattc 1500
cagttagggtc tggctcaggc attggacatt aaagtccata ctatgactcg tgaggttatg 1560
tctgacgagt tgactaaatt actggaaggt aatctgaaac cagccattga tatgatgggt 1620
gagtttaata ccactgggtc cttaccagaa aacgcagttg atgttctgaa tacagcatta 1680
ggagatagga agtcattcgt agcattgatg gctcttatgg agtattcccc ttacttagta 1740

Epicentre-00005 seqlist (Nov).txt

gcagaggata aatctgcatt tgtaactcca ctgtatgtag aagcagatgg tggtactaat 1800
 ggtccaatca atgccatgat gctaatagaca ggcggctctgt ttactcctga ctggattcgt 1860
 aatattgcca aaggggggctt gttcattgggt tctccaaata agaccatgaa tgagcatcgc 1920
 tctactgctg acaataatga tttatatcaa gcatccacta atgctttgat ggaatcggtg 1980
 ggtaagttac gtagtaacta tgcctctaata atgcctattc agtctcagat agacagtctt 2040
 ctttctctga tggatttggt tttaccggat attaactctg gtgagaatgg tgctttagaa 2100
 cttaaacgtg gtattgctaa gaaccactg actattacca tcttcgggtt tggtgctcgt 2160
 ggtattgcag gtaagctgggt tagttctggt actgatgcca tctatgagcg tatgtctgat 2220
 gtactgaaag ctctgctaa agacccaaat atctctgctg ctatggcaat gtttggtgaa 2280
 caagctgctt cagaagcaca tgctgaagaa cttcttgccc gtttcctgaa agatatggaa 2340
 aactgactt ctactgttcc tgttaaactg aaaggtgtac tggaactaca atccacaggt 2400
 acaggagcca aaggaaaaat caatcctaag acctatacca ttaagggcga gcaactgaag 2460
 gcacttcagg aaaatatgct gcacttcttt gtagaaccac tacgtaatgg tattactcag 2520
 actgtaggtg aaagtctgggt gtactctact gaacaattac agaaagctac tcagattcaa 2580
 tctgtagtgc tggaagatat gttcaaacag cgagtacaag agaagctggc agagaaggct 2640
 aaagacccaa catggaagaa aggtgatttc cttactcaga aagaactgaa tgatattcag 2700
 gcttctctga ataacttagc ccctatgatt gagactgggt ctcagacttt ctacattgct 2760
 ggttcagaaa atgcagaagt agcaaatcag gtattagcta ctaaccttga tgaccgtatg 2820
 cgtgtaccaa tgagtatcta tgctccagca caggccgggt tagcaggtat tccatttatg 2880
 actattggta ctggtgatgg catgatgatg caaactcttt ccactatgaa aggtgcacca 2940
 aagaataccc tcaaaatctt tgatgggtat aacattgggt tgaatgacat cactgatgcc 3000
 agtcgtaaag ctaatgaagc tgtttacact tcttggcagg gtaaccctat taagaatggt 3060
 tatgaatcat atgctaagtt catgaagaat gtagatttca gcaagctgtc ccctgaagca 3120
 ttggaagcaa ttggtaaact tgctctggaa tatgaccaac gtgagaatgc tactgtagat 3180

Epicentre-00005 seqlist (Nov).txt

gatattgcta acgctgcatc tctgattgaa cgtaacttac gtaatattgc actgggtgta 3240
gatattcgtc ataaggtgct ggataaggta aatctgtcca ttgaccagat ggctgctgta 3300
ggctgctcctt atcagaacaa cggttaagatt gacctcagca atatgacccc tgaacaacag 3360
gctgatgaac tgaataaact tttccgtgaa gagttagaag cccgtaaaca aaaagtcgct 3420
aaggctaggt aa 3432

<210> 8

<211> 1143

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 8

Met	Gly	Gly	Ser	His	His	His	His	His	His	Gly	Met	Ala	Ser	Met	Thr
1				5					10					15	
Gly	Gly	Gln	Gln	Met	Gly	Arg	Asp	Leu	Tyr	Asp	Asp	Asp	Asp	Lys	Asp
			20					25					30		
Pro	Ser	Ser	Arg	Ser	Glu	Ser	Thr	Val	Thr	Glu	Glu	Leu	Lys	Glu	Gly
			35				40					45			
Ile	Asp	Ala	Val	Tyr	Pro	Ser	Leu	Val	Gly	Thr	Ala	Asp	Ser	Lys	Ala
	50					55					60				
Glu	Gly	Ile	Lys	Asn	Tyr	Phe	Lys	Leu	Ser	Phe	Thr	Leu	Pro	Glu	Glu
	65				70					75				80	
Gln	Lys	Ser	Arg	Thr	Val	Gly	Ser	Glu	Ala	Pro	Leu	Lys	Asp	Val	Ala
				85					90					95	
Gln	Ala	Leu	Ser	Ser	Arg	Ala	Arg	Tyr	Glu	Leu	Phe	Thr	Glu	Lys	Glu
			100					105					110		
Thr	Ala	Asn	Pro	Ala	Phe	Asn	Gly	Glu	Val	Ile	Lys	Arg	Tyr	Lys	Glu
		115					120					125			
Leu	Met	Glu	His	Gly	Glu	Gly	Ile	Ala	Asp	Ile	Leu	Arg	Ser	Arg	Leu
	130					135					140				

Epicentre-00005 seqlist (Nov).txt

Ala	Lys	Phe	Leu	Asn	Thr	Lys	Asp	Val	Gly	Lys	Arg	Phe	Ala	Gln	Gly	145	150	155	160
Thr	Glu	Ala	Asn	Arg	Trp	Val	Gly	Gly	Lys	Leu	Leu	Asn	Ile	Val	Glu	165	170	175	
Gln	Asp	Gly	Asp	Thr	Phe	Lys	Tyr	Asn	Glu	Gln	Leu	Leu	Gln	Thr	Ala	180	185	190	
Val	Leu	Ala	Gly	Leu	Gln	Trp	Arg	Leu	Thr	Ala	Thr	Ser	Asn	Thr	Ala	195	200	205	
Ile	Lys	Asp	Ala	Lys	Asp	Val	Ala	Ala	Ile	Thr	Gly	Ile	Asp	Gln	Ala	210	215	220	
Leu	Leu	Pro	Glu	Gly	Leu	Val	Glu	Gln	Phe	Asp	Thr	Gly	Met	Thr	Leu	225	230	235	240
Thr	Glu	Ala	Val	Ser	Ser	Leu	Ala	Gln	Lys	Ile	Glu	Ser	Tyr	Trp	Gly	245	250	255	
Leu	Ser	Arg	Asn	Pro	Asn	Ala	Pro	Leu	Gly	Tyr	Thr	Lys	Gly	Ile	Pro	260	265	270	
Thr	Ala	Met	Ala	Ala	Glu	Ile	Leu	Ala	Ala	Phe	Val	Glu	Ser	Thr	Asp	275	280	285	
Val	Val	Glu	Asn	Ile	Val	Asp	Met	Ser	Glu	Ile	Asp	Pro	Asp	Asn	Lys	290	295	300	
Lys	Thr	Ile	Gly	Leu	Tyr	Thr	Ile	Thr	Glu	Leu	Asp	Ser	Phe	Asp	Pro	305	310	315	320
Ile	Asn	Ser	Phe	Pro	Thr	Ala	Ile	Glu	Glu	Ala	Val	Leu	Val	Asn	Pro	325	330	335	
Thr	Glu	Lys	Met	Phe	Phe	Gly	Asp	Asp	Ile	Pro	Pro	Val	Ala	Asn	Thr	340	345	350	
Gln	Leu	Arg	Asn	Pro	Ala	Val	Arg	Asn	Thr	Pro	Glu	Gln	Lys	Ala	Ala	355	360	365	
Leu	Lys	Ala	Glu	Gln	Ala	Thr	Glu	Phe	Tyr	Val	His	Thr	Pro	Met	Val	370	375	380	
Gln	Phe	Tyr	Glu	Thr	Leu	Gly	Lys	Asp	Arg	Ile	Leu	Glu	Leu	Met	Gly	385	390	395	400

Epicentre-00005 seqlist (Nov).txt

Ala	Gly	Thr	Leu	Asn	Lys	Glu	Leu	Leu	Asn	Asp	Asn	His	Ala	Lys	Ser	405	410	415
Leu	Glu	Gly	Lys	Asn	Arg	Ser	Val	Glu	Asp	Ser	Tyr	Asn	Gln	Leu	Phe	420	425	430
Ser	Val	Ile	Glu	Gln	Val	Arg	Ala	Gln	Ser	Glu	Asp	Ile	Ser	Thr	Val	435	440	445
Pro	Ile	His	Tyr	Ala	Tyr	Asn	Met	Thr	Arg	Val	Gly	Arg	Met	Gln	Met	450	455	460
Leu	Gly	Lys	Tyr	Asn	Pro	Gln	Ser	Ala	Lys	Leu	Val	Arg	Glu	Ala	Ile	465	470	475
Leu	Pro	Thr	Lys	Ala	Thr	Leu	Asp	Leu	Ser	Asn	Gln	Asn	Asn	Glu	Asp	485	490	495
Phe	Ser	Ala	Phe	Gln	Leu	Gly	Leu	Ala	Gln	Ala	Leu	Asp	Ile	Lys	Val	500	505	510
His	Thr	Met	Thr	Arg	Glu	Val	Met	Ser	Asp	Glu	Leu	Thr	Lys	Leu	Leu	515	520	525
Glu	Gly	Asn	Leu	Lys	Pro	Ala	Ile	Asp	Met	Met	Val	Glu	Phe	Asn	Thr	530	535	540
Thr	Gly	Ser	Leu	Pro	Glu	Asn	Ala	Val	Asp	Val	Leu	Asn	Thr	Ala	Leu	545	550	555
Gly	Asp	Arg	Lys	Ser	Phe	Val	Ala	Leu	Met	Ala	Leu	Met	Glu	Tyr	Ser	565	570	575
Arg	Tyr	Leu	Val	Ala	Glu	Asp	Lys	Ser	Ala	Phe	Val	Thr	Pro	Leu	Tyr	580	585	590
Val	Glu	Ala	Asp	Gly	Val	Thr	Asn	Gly	Pro	Ile	Asn	Ala	Met	Met	Leu	595	600	605
Met	Thr	Gly	Gly	Leu	Phe	Thr	Pro	Asp	Trp	Ile	Arg	Asn	Ile	Ala	Lys	610	615	620
Gly	Gly	Leu	Phe	Ile	Gly	Ser	Pro	Asn	Lys	Thr	Met	Asn	Glu	His	Arg	625	630	635
Ser	Thr	Ala	Asp	Asn	Asn	Asp	Leu	Tyr	Gln	Ala	Ser	Thr	Asn	Ala	Leu	645	650	655

Epicentre-00005 seqlist (Nov).txt

Met	Glu	Ser	Leu	Gly	Lys	Leu	Arg	Ser	Asn	Tyr	Ala	Ser	Asn	Met	Pro
			660					665					670		
Ile	Gln	Ser	Gln	Ile	Asp	Ser	Leu	Leu	Ser	Leu	Met	Asp	Leu	Phe	Leu
		675					680					685			
Pro	Asp	Ile	Asn	Leu	Gly	Glu	Asn	Gly	Ala	Leu	Glu	Leu	Lys	Arg	Gly
	690					695					700				
Ile	Ala	Lys	Asn	Pro	Leu	Thr	Ile	Thr	Ile	Phe	Gly	Ser	Gly	Ala	Arg
705					710					715					720
Gly	Ile	Ala	Gly	Lys	Leu	Val	Ser	Ser	Val	Thr	Asp	Ala	Ile	Tyr	Glu
				725					730					735	
Arg	Met	Ser	Asp	Val	Leu	Lys	Ala	Arg	Ala	Lys	Asp	Pro	Asn	Ile	Ser
			740					745					750		
Ala	Ala	Met	Ala	Met	Phe	Gly	Lys	Gln	Ala	Ala	Ser	Glu	Ala	His	Ala
		755					760					765			
Glu	Glu	Leu	Leu	Ala	Arg	Phe	Leu	Lys	Asp	Met	Glu	Thr	Leu	Thr	Ser
	770					775					780				
Thr	Val	Pro	Val	Lys	Arg	Lys	Gly	Val	Leu	Glu	Leu	Gln	Ser	Thr	Gly
785				790						795					800
Thr	Gly	Ala	Lys	Gly	Lys	Ile	Asn	Pro	Lys	Thr	Tyr	Thr	Ile	Lys	Gly
				805					810					815	
Glu	Gln	Leu	Lys	Ala	Leu	Gln	Glu	Asn	Met	Leu	His	Phe	Phe	Val	Glu
			820					825					830		
Pro	Leu	Arg	Asn	Gly	Ile	Thr	Gln	Thr	Val	Gly	Glu	Ser	Leu	Val	Tyr
		835					840					845			
Ser	Thr	Glu	Gln	Leu	Gln	Lys	Ala	Thr	Gln	Ile	Gln	Ser	Val	Val	Leu
	850					855					860				
Glu	Asp	Met	Phe	Lys	Gln	Arg	Val	Gln	Glu	Lys	Leu	Ala	Glu	Lys	Ala
865					870					875					880
Lys	Asp	Pro	Thr	Trp	Lys	Lys	Gly	Asp	Phe	Leu	Thr	Gln	Lys	Glu	Leu
				885					890					895	
Asn	Asp	Ile	Gln	Ala	Ser	Leu	Asn	Asn	Leu	Ala	Pro	Met	Ile	Glu	Thr
			900					905					910		

Epicentre-00005 seqlist (Nov).txt

[illegible]

Epicentre-00005 seqlist (Nov).txt

<211> 69
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

<400> 9
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cggagcttc 69

<210> 10
<211> 69
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

<400> 10
tgggagagaa aaggттаага tttgatagag gattggatgg attagttgat ggaaaагааg 60
cggagcttc 69

<210> 11
<211> 69
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

<400> 11
tcсctgtctt ttggttttgt ttctttctg gtttggttgc ttttcttctt ccaaаагааg 60
cggagcttc 69

<210> 12

Epicentre-00005 seqlist (Nov).txt

<211> 69
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

<400> 12
tcccacacaa aaccttaaca ttccatacac cattccatcc attacttcat ccaaagaag 60
cggagcttc 69

<210> 13
<211> 69
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

<400> 13
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cggagcttc 69

<210> 14
<211> 10617
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

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tttgatagac tggctgggtt cgcagacagc gtaaccaatg caaagcaagt tgacgtctct 180
actgcaaccg cccagaagaa agctgaacaa ggtgtcacta ctctcttgt ttctctgtat 240

Epicentre-00005 seqlist (Nov).txt

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Epicentre-00005 seqlist (Nov).txt

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Epicentre-00005 seqlist (Nov).txt

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Epicentre-00005 seqlist (Nov).txt

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Epicentre-00005 seqlist (Nov).txt

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Epicentre-00005 seqlist (Nov).txt

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Epicentre-00005 seqlist (Nov).txt

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Epicentre-00005 seqlist (Nov).txt

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7

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<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 15

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Pro Ser Ser Arg Ser Met Ser Val Phe Asp Arg Leu Ala Gly Phe Ala
      35                      40                      45

Asp Ser Val Thr Asn Ala Lys Gln Val Asp Val Ser Thr Ala Thr Ala
      50                      55                      60

Gln Lys Lys Ala Glu Gln Gly Val Thr Thr Pro Leu Val Ser Pro Asp
      65                      70                      75                      80

Ala Ala Tyr Gln Met Gln Ala Ala Arg Thr Gly Asn Val Gly Ala Asn
      85                      90                      95

Ala Phe Glu Pro Gly Thr Val Gln Ser Asp Phe Met Asn Leu Thr Pro
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Met Gln Ile Met Asn Lys Tyr Gly Val Glu Gln Gly Leu Gln Leu Ile
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Epicentre-00005 seqlist (Nov).txt

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Epicentre-00005 seqlist (Nov).txt

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Asn	Tyr	Gln	Glu	Ala	Pro	Glu	His	Ile	Arg	Asn	Ala	Val	Ala	Gly	Ser
		595					600					605			
Thr	Asp	Gln	Val	Gln	Val	Ile	Gln	Lys	Leu	Ala	Asp	Leu	Val	Asn	Thr
	610					615					620				
Leu	Asp	Glu	Ser	Asn	Pro	Gln	Ala	Leu	Met	Glu	Ala	Ala	Ser	Tyr	Met
625					630					635					640
Tyr	Asp	Ala	Val	Ser	Glu	Phe	Glu	Gln	Phe	Ile	Asn	Arg	Asp	Pro	Ala

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645 650 655

Ala	Leu	Asp	Ser	Ile	Pro	Lys	Asp	Ser	Pro	Ala	Ile	Glu	Leu	Leu	Asn
			660					665					670		
Arg	Tyr	Thr	Asn	Leu	Thr	Ala	Asn	Ile	Gln	Asn	Thr	Pro	Lys	Val	Ile
		675					680					685			
Gly	Ala	Leu	Asn	Val	Ile	Asn	Arg	Met	Ile	Asn	Glu	Ser	Ala	Gln	Asn
	690					695					700				
Gly	Ser	Leu	Asn	Val	Thr	Glu	Glu	Ser	Ser	Pro	Gln	Glu	Met	Gln	Asn
705					710					715					720
Val	Ala	Leu	Ala	Ala	Glu	Val	Ala	Pro	Glu	Lys	Leu	Asn	Pro	Glu	Ser
				725					730					735	
Val	Asn	Val	Val	Leu	Lys	His	Ala	Ala	Asp	Gly	Arg	Ile	Lys	Leu	Asn
			740					745					750		
Asn	Arg	Gln	Ile	Ala	Ala	Leu	Gln	Asn	Ala	Ala	Ala	Ile	Leu	Lys	Gly
		755					760					765			
Ala	Arg	Glu	Tyr	Asp	Ala	Glu	Ala	Ala	Arg	Leu	Gly	Leu	Arg	Pro	Gln
	770					775					780				
Asp	Ile	Val	Ser	Lys	Gln	Ile	Lys	Thr	Asp	Glu	Ser	Arg	Thr	Gln	Glu
785					790					795					800
Gly	Gln	Tyr	Ser	Ala	Leu	Gln	His	Ala	Asn	Arg	Ile	Arg	Ser	Ala	Tyr
				805					810					815	
Asn	Ser	Gly	Asn	Phe	Glu	Leu	Ala	Ser	Ala	Tyr	Leu	Asn	Asp	Phe	Met
			820					825					830		
Gln	Phe	Ala	Gln	His	Met	Gln	Asn	Lys	Val	Gly	Ala	Leu	Asn	Glu	His
		835					840					845			
Leu	Val	Thr	Gly	Asn	Ala	Asp	Lys	Asn	Lys	Ser	Val	His	Tyr	Gln	Ala
	850					855					860				
Leu	Thr	Ala	Asp	Arg	Glu	Trp	Val	Arg	Ser	Arg	Thr	Gly	Leu	Gly	Val
865					870					875					880
Asn	Pro	Tyr	Asp	Thr	Lys	Ser	Val	Lys	Phe	Ala	Gln	Gln	Val	Ala	Leu
				885					890					895	
Glu	Ala	Lys	Thr	Val	Ala	Asp	Ile	Ala	Asn	Ala	Leu	Ala	Ser	Ala	Tyr

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900								905						910	
Pro	Glu	Leu	Lys	Val	Ser	His	Ile	Lys	Val	Thr	Pro	Leu	Asp	Ser	Arg
		915					920					925			
Leu	Asn	Ala	Pro	Ala	Ala	Glu	Val	Val	Lys	Ala	Phe	Arg	Gln	Gly	Asn
	930					935					940				
Arg	Asp	Val	Ala	Ser	Ser	Gln	Pro	Lys	Ala	Asp	Ser	Val	Asn	Gln	Val
945					950					955					960
Lys	Glu	Thr	Pro	Val	Thr	Lys	Gln	Glu	Pro	Val	Thr	Ser	Thr	Val	Gln
				965					970					975	
Thr	Lys	Thr	Pro	Val	Ser	Glu	Ser	Val	Lys	Thr	Glu	Pro	Thr	Thr	Lys
			980					985					990		
Glu	Ser	Ser	Pro	Gln	Ala	Ile	Lys	Glu	Pro	Val	Asn	Gln	Ser	Glu	Lys
		995					1000					1005			
Gln	Asp	Val	Asn	Leu	Thr	Asn	Glu	Asp	Asn	Ile	Lys	Gln	Pro	Thr	Glu
	1010					1015					1020				
Ser	Val	Lys	Glu	Thr	Glu	Thr	Ser	Thr	Lys	Glu	Ser	Thr	Val	Thr	Glu
1025					1030					1035					1040
Glu	Leu	Lys	Glu	Gly	Ile	Asp	Ala	Val	Tyr	Pro	Ser	Leu	Val	Gly	Thr
				1045					1050					1055	
Ala	Asp	Ser	Lys	Ala	Glu	Gly	Ile	Lys	Asn	Tyr	Phe	Lys	Leu	Ser	Phe
			1060					1065					1070		
Thr	Leu	Pro	Glu	Glu	Gln	Lys	Ser	Arg	Thr	Val	Gly	Ser	Glu	Ala	Pro
		1075					1080					1085			
Leu	Lys	Asp	Val	Ala	Gln	Ala	Leu	Ser	Ser	Arg	Ala	Arg	Tyr	Glu	Leu
	1090					1095					1100				
Phe	Thr	Glu	Lys	Glu	Thr	Ala	Asn	Pro	Ala	Phe	Asn	Gly	Glu	Val	Ile
1105					1110					1115				1120	
Lys	Arg	Tyr	Lys	Glu	Leu	Met	Glu	His	Gly	Glu	Gly	Ile	Ala	Asp	Ile
				1125					1130					1135	
Leu	Arg	Ser	Arg	Leu	Ala	Lys	Phe	Leu	Asn	Thr	Lys	Asp	Val	Gly	Lys
			1140					1145					1150		
Arg	Phe	Ala	Gln	Gly	Thr	Glu	Ala	Asn	Arg	Trp	Val	Gly	Gly	Lys	Leu

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1155	1160	1165
Leu Asn Ile Val Glu Gln Asp Gly Asp Thr Phe Lys Tyr Asn Glu Gln		
1170	1175	1180
Leu Leu Gln Thr Ala Val Leu Ala Gly Leu Gln Trp Arg Leu Thr Ala		
1185	1190	1195 1200
Thr Ser Asn Thr Ala Ile Lys Asp Ala Lys Asp Val Ala Ala Ile Thr		
	1205 1210	1215
Gly Ile Asp Gln Ala Leu Leu Pro Glu Gly Leu Val Glu Gln Phe Asp		
	1220 1225	1230
Thr Gly Met Thr Leu Thr Glu Ala Val Ser Ser Leu Ala Gln Lys Ile		
	1235 1240	1245
Glu Ser Tyr Trp Gly Leu Ser Arg Asn Pro Asn Ala Pro Leu Gly Tyr		
	1250 1255	1260
Thr Lys Gly Ile Pro Thr Ala Met Ala Ala Glu Ile Leu Ala Ala Phe		
	1265 1270	1275 1280
Val Glu Ser Thr Asp Val Val Glu Asn Ile Val Asp Met Ser Glu Ile		
	1285 1290	1295
Asp Pro Asp Asn Lys Lys Thr Ile Gly Leu Tyr Thr Ile Thr Glu Leu		
	1300 1305	1310
Asp Ser Phe Asp Pro Ile Asn Ser Phe Pro Thr Ala Ile Glu Glu Ala		
	1315 1320	1325
Val Leu Val Asn Pro Thr Glu Lys Met Phe Phe Gly Asp Asp Ile Pro		
	1330 1335	1340
Pro Val Ala Asn Thr Gln Leu Arg Asn Pro Ala Val Arg Asn Thr Pro		
	1345 1350	1355 1360
Glu Gln Lys Ala Ala Leu Lys Ala Glu Gln Ala Thr Glu Phe Tyr Val		
	1365 1370	1375
His Thr Pro Met Val Gln Phe Tyr Glu Thr Leu Gly Lys Asp Arg Ile		
	1380 1385	1390
Leu Glu Leu Met Gly Ala Gly Thr Leu Asn Lys Glu Leu Leu Asn Asp		
	1395 1400	1405
Asn His Ala Lys Ser Leu Glu Gly Lys Asn Arg Ser Val Glu Asp Ser		

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1410	1415	1420
Tyr Asn Gln Leu Phe Ser Val Ile Glu Gln Val Arg Ala Gln Ser Glu		
1425	1430	1435 1440
Asp Ile Ser Thr Val Pro Ile His Tyr Ala Tyr Asn Met Thr Arg Val		
	1445	1450 1455
Gly Arg Met Gln Met Leu Gly Lys Tyr Asn Pro Gln Ser Ala Lys Leu		
	1460	1465 1470
Val Arg Glu Ala Ile Leu Pro Thr Lys Ala Thr Leu Asp Leu Ser Asn		
	1475	1480 1485
Gln Asn Asn Glu Asp Phe Ser Ala Phe Gln Leu Gly Leu Ala Gln Ala		
	1490	1495 1500
Leu Asp Ile Lys Val His Thr Met Thr Arg Glu Val Met Ser Asp Glu		
	1505	1510 1515 1520
Leu Thr Lys Leu Leu Glu Gly Asn Leu Lys Pro Ala Ile Asp Met Met		
	1525	1530 1535
Val Glu Phe Asn Thr Thr Gly Ser Leu Pro Glu Asn Ala Val Asp Val		
	1540	1545 1550
Leu Asn Thr Ala Leu Gly Asp Arg Lys Ser Phe Val Ala Leu Met Ala		
	1555	1560 1565
Leu Met Glu Tyr Ser Arg Tyr Leu Val Ala Glu Asp Lys Ser Ala Phe		
	1570	1575 1580
Val Thr Pro Leu Tyr Val Glu Ala Asp Gly Val Thr Asn Gly Pro Ile		
	1585	1590 1595 1600
Asn Ala Met Met Leu Met Thr Gly Gly Leu Phe Thr Pro Asp Trp Ile		
	1605	1610 1615
Arg Asn Ile Ala Lys Gly Gly Leu Phe Ile Gly Ser Pro Asn Lys Thr		
	1620	1625 1630
Met Asn Glu His Arg Ser Thr Ala Asp Asn Asn Asp Leu Tyr Gln Ala		
	1635	1640 1645
Ser Thr Asn Ala Leu Met Glu Ser Leu Gly Lys Leu Arg Ser Asn Tyr		
	1650	1655 1660
Ala Ser Asn Met Pro Ile Gln Ser Gln Ile Asp Ser Leu Leu Ser Leu		

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1665	1670	1675	1680
Met Asp Leu Phe Leu Pro Asp Ile Asn Leu Gly Glu Asn Gly Ala Leu	1685	1690	1695
Glu Leu Lys Arg Gly Ile Ala Lys Asn Pro Leu Thr Ile Thr Ile Tyr	1700	1705	1710
Gly Ser Gly Ala Arg Gly Ile Ala Gly Lys Leu Val Ser Ser Val Thr	1715	1720	1725
Asp Ala Ile Tyr Glu Arg Met Ser Asp Val Leu Lys Ala Arg Ala Lys	1730	1735	1740
Asp Pro Asn Ile Ser Ala Ala Met Ala Met Phe Gly Lys Gln Ala Ala	1745	1750	1755
Ser Glu Ala His Ala Glu Glu Leu Leu Ala Arg Phe Leu Lys Asp Met	1765	1770	1775
Glu Thr Leu Thr Ser Thr Val Pro Val Lys Arg Lys Gly Val Leu Glu	1780	1785	1790
Leu Gln Ser Thr Gly Thr Gly Ala Lys Gly Lys Ile Asn Pro Lys Thr	1795	1800	1805
Tyr Thr Ile Lys Gly Glu Gln Leu Lys Ala Leu Gln Glu Asn Met Leu	1810	1815	1820
His Phe Phe Val Glu Pro Leu Arg Asn Gly Ile Thr Gln Thr Val Gly	1825	1830	1835
Glu Ser Leu Val Tyr Ser Thr Glu Gln Leu Gln Lys Ala Thr Gln Ile	1845	1850	1855
Gln Ser Val Val Leu Glu Asp Met Phe Lys Gln Arg Val Gln Glu Lys	1860	1865	1870
Leu Ala Glu Lys Ala Lys Asp Pro Thr Trp Lys Lys Gly Asp Phe Leu	1875	1880	1885
Thr Gln Lys Glu Leu Asn Asp Ile Gln Ala Ser Leu Asn Asn Leu Ala	1890	1895	1900
Pro Met Ile Glu Thr Gly Ser Gln Thr Phe Tyr Ile Ala Gly Ser Glu	1905	1910	1915
Asn Ala Glu Val Ala Asn Gln Val Leu Ala Thr Asn Leu Asp Asp Arg			

1925 1930 1935

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2180		2185		2190
Ala Thr Leu Ala Glu Ile Gln Lys Ser Leu Ala Ala Lys Asp Tyr Lys	2195	2200	2205	
Ile Ile Tyr Gly Thr Pro Thr Gln Val Ala Glu Tyr Ala Arg Gln Lys	2210	2215	2220	
Asn Val Thr Glu Leu Thr Ser Gln Glu Met Glu Glu Ala Gln Ala Gly	2225	2230	2235	2240
Asn Ile Tyr Gly Trp Thr Asn Phe Asp Asp Lys Thr Ile Tyr Leu Val	2245	2250	2255	
Ser Pro Ser Met Glu Thr Leu Ile His Glu Leu Val His Ala Ser Thr	2260	2265	2270	
Phe Glu Glu Val Tyr Ser Phe Tyr Gln Gly Asn Glu Val Ser Pro Thr	2275	2280	2285	
Ser Lys Gln Ala Ile Glu Asn Leu Glu Gly Leu Met Glu Gln Phe Arg	2290	2295	2300	
Ser Leu Asp Ile Ser Lys Asp Ser Pro Glu Met Arg Glu Ala Tyr Ala	2305	2310	2315	2320
Asp Ala Ile Ala Thr Ile Glu Gly His Leu Ser Asn Gly Phe Val Asp	2325	2330	2335	
Pro Ala Ile Ser Lys Ala Ala Ala Leu Asn Glu Phe Met Ala Trp Gly	2340	2345	2350	
Leu Ala Asn Arg Ala Leu Ala Ala Lys Gln Lys Arg Thr Ser Ser Leu	2355	2360	2365	
Val Gln Met Val Lys Asp Val Tyr Gln Ala Ile Lys Lys Leu Ile Trp	2370	2375	2380	
Gly Arg Lys Gln Ala Pro Ala Leu Gly Glu Asp Met Phe Ser Asn Leu	2385	2390	2395	2400
Leu Phe Asn Ser Ala Ile Leu Met Arg Ser Gln Pro Thr Thr Gln Ala	2405	2410	2415	
Val Ala Lys Asp Gly Thr Leu Phe His Ser Lys Ala Tyr Gly Asn Asn	2420	2425	2430	
Glu Arg Leu Ser Gln Leu Asn Gln Thr Phe Asp Lys Leu Val Thr Asp				

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2435	2440	2445
Tyr Leu Arg Thr Asp Pro Val Thr Glu Val Glu Arg Arg Gly Asn Val		
2450	2455	2460
Ala Asn Ala Leu Met Ser Ala Thr Arg Leu Val Arg Asp Val Gln Ser		
2465	2470	2475 2480
His Gly Phe Asn Met Thr Ala Gln Glu Gln Ser Val Phe Gln Met Val		
2485	2490	2495
Thr Ala Ala Leu Ala Thr Glu Ala Ala Ile Asp Pro His Ala Met Ala		
2500	2505	2510
Arg Ala Gln Glu Leu Tyr Thr His Val Met Lys His Leu Thr Val Glu		
2515	2520	2525
His Phe Met Ala Asp Pro Asp Ser Thr Asn Pro Ala Asp Arg Tyr Tyr		
2530	2535	2540
Ala Gln Gln Lys Tyr Asp Thr Ile Ser Gly Ala Asn Leu Val Glu Val		
2545	2550	2555 2560
Asp Ala Lys Gly Arg Thr Ser Leu Leu Pro Thr Phe Leu Gly Leu Ala		
2565	2570	2575
Met Val Asn Glu Glu Leu Arg Ser Ile Ile Lys Glu Met Pro Val Pro		
2580	2585	2590
Lys Ala Asp Lys Lys Leu Gly Asn Asp Ile Asp Thr Leu Leu Thr Asn		
2595	2600	2605
Ala Gly Thr Gln Val Met Glu Ser Leu Asn Arg Arg Met Ala Gly Asp		
2610	2615	2620
Gln Lys Ala Thr Asn Val Gln Asp Ser Ile Asp Ala Leu Ser Glu Thr		
2625	2630	2635 2640
Ile Met Ala Ala Ala Leu Lys Arg Glu Ser Phe Tyr Asp Ala Val Ala		
2645	2650	2655
Thr Pro Thr Gly Asn Phe Ile Asp Arg Ala Asn Gln Tyr Val Thr Asp		
2660	2665	2670
Ser Ile Glu Arg Leu Ser Glu Thr Val Ile Glu Lys Ala Asp Lys Val		
2675	2680	2685
Ile Ala Asn Pro Ser Asn Ile Ala Ala Lys Gly Val Ala His Leu Ala		

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2690

2695

2700

Lys Leu Thr Ala Ala Ile Ala Ser Glu Lys Gln Gly Glu Ile Val Ala
2705 2710 2715 2720

Gln Gly Val Met Thr Ala Met Asn Gln Gly Lys Val Trp Gln Pro Phe
2725 2730 2735

His Asp Leu Val Asn Asp Ile Val Gly Arg Thr Lys Thr Asn Ala Asn
2740 2745 2750

Val Tyr Asp Leu Ile Lys Leu Val Lys Ser Gln Ile Ser Gln Asp Arg
2755 2760 2765

Gln Gln Phe Arg Glu His Leu Pro Thr Val Ile Ala Gly Lys Phe Ser
2770 2775 2780

Arg Lys Leu Thr Asp Thr Glu Trp Ser Ala Met His Thr Gly Leu Gly
2785 2790 2795 2800

Lys Thr Asp Leu Ala Val Leu Arg Glu Thr Met Ser Met Ala Glu Ile
2805 2810 2815

Arg Asp Leu Leu Ser Ser Ser Lys Lys Val Lys Asp Glu Ile Ser Thr
2820 2825 2830

Leu Glu Lys Glu Ile Gln Asn Gln Ala Gly Arg Asn Trp Asn Leu Val
2835 2840 2845

Gln Lys Lys Ser Lys Gln Leu Ala Gln Tyr Met Ile Met Gly Glu Val
2850 2855 2860

Gly Asn Asn Leu Leu Arg Asn Ala His Ala Ile Ser Arg Leu Leu Gly
2865 2870 2875 2880

Glu Arg Ile Thr Asn Gly Pro Val Ala Asp Val Ala Ala Ile Asp Lys
2885 2890 2895

Leu Ile Thr Leu Tyr Ser Leu Glu Leu Met Asn Lys Ser Asp Arg Asp
2900 2905 2910

Leu Leu Ser Glu Leu Ala Gln Ser Glu Val Glu Gly Met Glu Phe Ser
2915 2920 2925

Ile Ala Tyr Met Val Gly Gln Arg Thr Glu Glu Met Arg Lys Ala Lys
2930 2935 2940

Gly Asp Asn Arg Thr Leu Leu Asn His Phe Lys Gly Tyr Ile Pro Val

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2945	2950	2955	2960
Glu Asn Gln Gln Gly Val Asn Leu Ile Ile Ala Asp Asp Lys Glu Phe	2965	2970	2975
Ala Lys Leu Asn Ser Gln Ser Phe Thr Arg Ile Gly Thr Tyr Gln Gly	2980	2985	2990
Ser Thr Gly Phe Arg Thr Gly Ser Lys Gly Tyr Tyr Phe Ser Pro Val	2995	3000	3005
Ala Ala Arg Ala Pro Tyr Ser Gln Gly Ile Leu Gln Asn Val Arg Asn	3010	3015	3020
Thr Ala Gly Gly Val Asp Ile Gly Thr Gly Phe Thr Leu Gly Thr Met	3025	3030	3040
Val Ala Gly Arg Ile Thr Asp Lys Pro Thr Val Glu Arg Ile Thr Lys	3045	3050	3055
Ala Leu Ala Lys Gly Glu Arg Gly Arg Glu Pro Leu Met Pro Ile Tyr	3060	3065	3070
Asn Ser Lys Gly Gln Val Val Ala Tyr Glu Gln Ser Val Asp Pro Asn	3075	3080	3085
Met Leu Lys His Leu Asn Gln Asp Asn His Phe Ala Lys Met Val Gly	3090	3095	3100
Val Trp Arg Gly Arg Gln Val Glu Glu Ala Lys Ala Gln Arg Phe Asn	3105	3110	3120
Asp Ile Leu Ile Glu Gln Leu His Ala Met Tyr Glu Lys Asp Ile Lys	3125	3130	3135
Asp Ser Ser Ala Asn Lys Ser Gln Tyr Val Asn Leu Leu Gly Lys Ile	3140	3145	3150
Asp Asp Pro Val Leu Ala Asp Ala Ile Asn Leu Met Asn Ile Glu Thr	3155	3160	3165
Arg His Lys Ala Glu Glu Leu Phe Gly Lys Asp Glu Leu Trp Val Arg	3170	3175	3180
Arg Asp Met Leu Asn Asp Ala Leu Gly Tyr Arg Ala Ala Ser Ile Gly	3185	3190	3200
Asp Val Trp Thr Gly Asn Ser Arg Trp Ser Pro Ser Thr Leu Asp Thr			

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3205 3210 3215

Val Lys Lys Met Phe Leu Gly Ala Phe Gly Asn Lys Ala Tyr His Val
3220 3225 3230

Val Met Asn Ala Glu Asn Thr Ile Gln Asn Leu Val Lys Asp Ala Lys
3235 3240 3245

Thr Val Ile Val Val Lys Ser Val Val Val Pro Ala Val Asn Phe Leu
3250 3255 3260

Ala Asn Ile Tyr Gln Met Ile Gly Arg Gly Val Pro Val Lys Asp Ile
3265 3270 3275 3280

Ala Val Asn Ile Pro Arg Lys Thr Ser Glu Ile Asn Gln Tyr Ile Lys
3285 3290 3295

Ser Arg Leu Arg Gln Ile Asp Ala Glu Ala Glu Leu Arg Ala Ala Glu
3300 3305 3310

Gly Asn Pro Asn Leu Val Arg Lys Leu Lys Thr Glu Ile Gln Ser Ile
3315 3320 3325

Thr Asp Ser His Arg Arg Met Ser Ile Trp Pro Leu Ile Glu Ala Gly
3330 3335 3340

Glu Phe Ser Ser Ile Ala Asp Ala Gly Ile Ser Arg Asp Asp Leu Leu
3345 3350 3355 3360

Val Ala Glu Gly Lys Ile His Glu Tyr Met Glu Lys Leu Ala Asn Lys
3365 3370 3375

Leu Pro Glu Lys Val Arg Asn Ala Gly Arg Tyr Ala Leu Ile Ala Lys
3380 3385 3390

Asp Thr Ala Leu Phe Gln Gly Ile Gln Lys Thr Val Glu Tyr Ser Asp
3395 3400 3405

Phe Ile Ala Lys Ala Ile Ile Tyr Asp Asp Leu Val Lys Arg Lys Lys
3410 3415 3420

Lys Ser Ser Ser Glu Ala Leu Gly Gln Val Thr Glu Glu Phe Ile Asn
3425 3430 3435 3440

Tyr Asp Arg Leu Pro Gly Arg Phe Arg Gly Tyr Met Glu Ser Met Gly
3445 3450 3455

Leu Met Trp Phe Tyr Asn Phe Lys Ile Arg Ser Ile Lys Val Ala Met

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3460 3465 3470

Ser Met Ile Arg Asn Asn Pro Val His Ser Leu Ile Ala Thr Val Val
3475 3480 3485

Pro Ala Pro Thr Met Phe Gly Asn Val Gly Leu Pro Ile Gln Asp Asn
3490 3495 3500

Met Leu Thr Met Leu Ala Glu Gly Arg Leu Asp Tyr Ser Leu Gly Phe
3505 3510 3515 3520

Gly Gln Gly Leu Arg Ala Pro Thr Leu Asn Pro Trp Phe Asn Leu Thr
3525 3530 3535

His

<210> 16
<211> 32
<212> DNA
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<220>
<223> Description of Artificial Sequence: Synthetic
Primer

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32

<210> 17
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<212> DNA
<213> Artificial Sequence

<220>
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Primer

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37

<210> 18
<211> 23

Epicentre-00005 seqlist (Nov).txt

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

<400> 18
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<210> 19
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

<400> 19
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<210> 20
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

<400> 20
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<210> 21
<211> 24
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<213> Artificial Sequence

<220>
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Primer

Epicentre-00005 seqlist (Nov).txt

<400> 21
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<210> 22
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

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<210> 23
<211> 33
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

<400> 23
agaagggggc tactaagccc tcttcttatt ttt 33

<210> 24
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<220>
<223> Description of Artificial Sequence: Synthetic
Primer

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Epicentre-00005 seqlist (Nov).txt

<210> 25
<211> 35
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

<400> 25
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<210> 26
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

<400> 26
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<210> 27
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
Primer

<400> 27
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<210> 28
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<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
Primer

<400> 28

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22

<210> 29

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
Primer

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20